
Sequence Listing could not be accepted.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Keisha Douglas

Timestamp: [year=2008; month=9; day=9; hr=16; min=48; sec=44; ms=79;]

Reviewer Comments: SEQUENCE LISTING

<110> DeveloGen AG f?wicklungsbiologische Forschung

<120> Use of a DG001 secreted protein product for preventing and treating pancreatic diseases and/or obesity and/or metabolic syndrome

<130> 31160PWO GE

Per the above sample, foreign accents are non-ascii characters which can not be processed.

Validated By CRFValidator v 1.0.3

Application No: 10560769 Version No: 1.0

Input Set:

Output Set:

Started: 2008-08-07 16:29:11.165

Finished: 2008-08-07 16:29:11.514

Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 349 ms

Total Warnings: 5

Total Errors: 0

No. of SeqIDs Defined: 5

Actual SeqID Count: 5

Error code		Error Description
W	402	Undefined organism found in <213> in SEQ ID (1)
W	402	Undefined organism found in <213> in SEQ ID (2)
W	213	Artificial or Unknown found in <213> in SEQ ID (3)
W	213	Artificial or Unknown found in <213> in SEQ ID (4)
W	213	Artificial or Unknown found in <213> in SEO ID (5)

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<211> 1300
<212> DNA
<213> human
<220>
<221> gene
<222> (1)..(1300)
<223> nucleic acid sequence encoding the human DG001
     protein
<400> 1
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ccattteect teegtteect eeetgteagg gegtaattga gteaaaggea ggateaggtt 120
ccccqccttc caqtccaaaa atcccqccaa qaqaqcccca qaqcaqaqqa aaatccaaaq 180
tggagagagg ggaagaaaga gaccagtgag tcatccgtcc agaaggcggg gagagcagca 240
geggeecaag eaggagetge agegageegg gtaeetggae teageggtag eaacetegee 300
ccttgcaaca aaggcagact gagcgccaga gaggacgttt ccaactcaaa aatgcaggct 360
caacagtacc agcagcagcg tcgaaaattt gcagctgcct tcttggcatt cattttcata 420
ctggcagctg tggatactgc tgaagcaggg aagaaagaga aaccagaaaa aaaagtgaag 480
aagtctgact gtggagaatg gcagtggagt gtgtgtgtgc ccaccagtgg agactgtggg 540
ctgggcacac gggagggcac tcggactgga gctgagtgca agcaaaccat gaagacccag 600
agatgtaaga tcccctgcaa ctggaagaag caatttggcg cggagtgcaa ataccagttc 660
caggcctggg gagaatgtga cctgaacaca gccctgaaga ccagaactgg aagtctgaag 720
cgagccctgc acaatgccga atgccagaag actgtcacca tctccaagcc ctgtggcaaa 780
ctgaccaagc ccaaacctca agcagaatct aagaagaaga aaaaggaagg caagaaacag 840
gagaagatgc tggattaaaa gatgtcacct gtggaacata aaaaggacat cagcaaacag 900
gatcagttaa ctattgcatt tatatgtacc gtaggctttg tattcaaaaa ttatctatag 960
ctaagtacac aataagcaaa aacaaaaaga aaagaaaatt tttgtagtag cgttttttaa 1020
atgtatacta tagtaccagt aggggcttat aataaaggac tgtaatctta tttaggaagt 1080
tgacttatag tacatgataa atgatagaca attgaggtaa gttttttgaa attatgtgac 1140
attttacatt aaatttttt tacatttttt gggcagcaat ttaaatgtta tgactatgta 1200
aactacttct cttgttaggt aatttttttc acctagattt ttttcccaat tgagaaaaat 1260
1300
<210> 2
<211> 168
<212> PRT
<213> human
<220>
<223> amino acid sequence of human DG001 protein
<400> 2
Met Gln Ala Gln Gln Tyr Gln Gln Gln Arg Arg Lys Phe Ala Ala Ala
 1
                                    10
Phe Leu Ala Phe Ile Phe Ile Leu Ala Ala Val Asp Thr Ala Glu Ala
                                25
            2.0
Gly Lys Lys Glu Lys Pro Glu Lys Lys Val Lys Lys Ser Asp Cys Gly
                            40
Glu Trp Gln Trp Ser Val Cys Val Pro Thr Ser Gly Asp Cys Gly Leu
    50
                        55
                                            60
```

<210> 1

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Gly Thr Arg Glu Gly Thr Arg Thr Gly Ala Glu Cys Lys Gln Thr Met
                     70
                                        75
Lys Thr Gln Arg Cys Lys Ile Pro Cys Asn Trp Lys Lys Gln Phe Gly
                85
                                    90
Ala Glu Cys Lys Tyr Gln Phe Gln Ala Trp Gly Glu Cys Asp Leu Asn
           100
                             105
Thr Ala Leu Lys Thr Arg Thr Gly Ser Leu Lys Arg Ala Leu His Asn
       115
                          120
                                              125
Ala Glu Cys Gln Lys Thr Val Thr Ile Ser Lys Pro Cys Gly Lys Leu
                    135
                                          140
Thr Lys Pro Lys Pro Gln Ala Glu Ser Lys Lys Lys Lys Glu Gly
145
                   150
                                       155
Lys Lys Gln Glu Lys Met Leu Asp
               165
<210> 3
<211> 22
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: primer 5`-3
<220>
<223> mouse DG001 forward primer
<400> 3
caagtaccag ttccaggctt gg
                                                                  22
<210> 4
<211> 17
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: primer 5`-3
<223> mouse DG001 reverse primer
<400> 4
                                                                 17
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<210> 5
<211> 30
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<212> DNA

<213> Artificial Sequence

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<220>
<223> Description of Artificial Sequence: probe
<220>
<223> mouse DG001 Taqman probe
<400> 5
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